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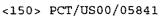
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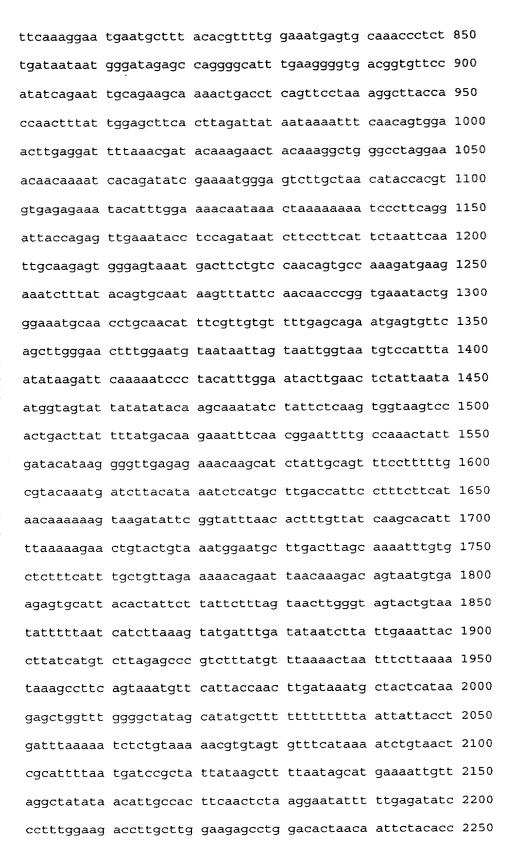
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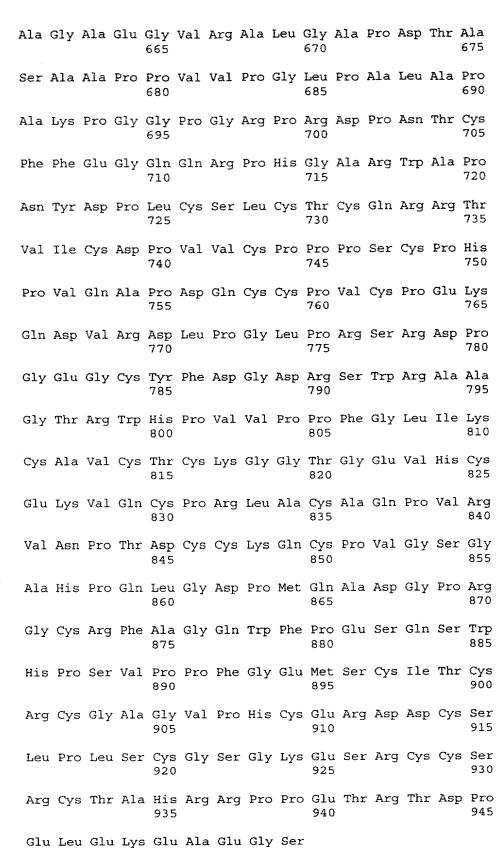
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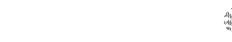
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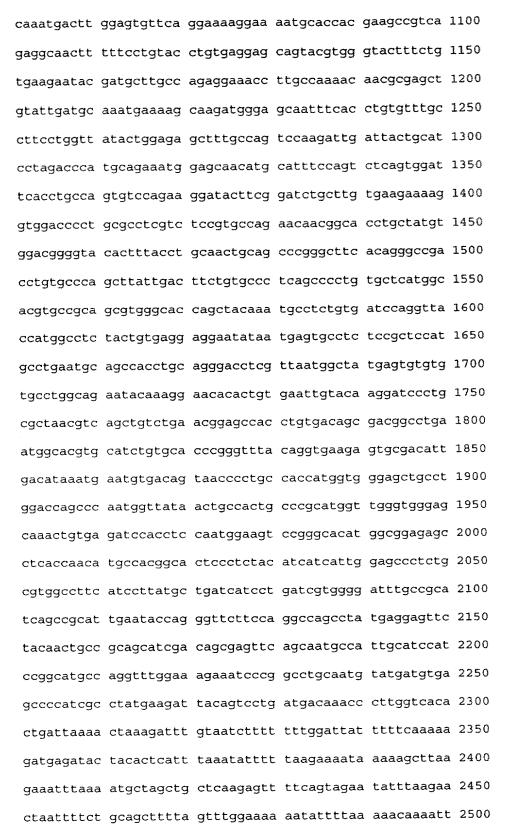
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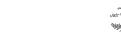
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Arg	Gln	Leu	Gln	Pro 155	Val	Pro	Ala	Thr	Gln 160	Glu	Pro	Asp	Lys	Ile 165
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Ala	Gly	Gly	Arg	Leu 215	Val	Ser	Phe	Glu	Val 220	Pro	Gln	Asn	Thr	Ser 225
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Lys	Val	Thr	Ala	Thr 245		Phe	Gln	Gln	Cys 250	Ser	Leu	Ile	Asp	Gly 255
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Glu	Glu	Met	Leu	Ala 275		Gly	Asn	Asn	His 280		Ile	Gly	Phe	Val 285
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Val	Val	Lys	Val	Ser 305		Cys	val	Pro	Gly 310		Ser	His	Ala	Asn 315
Asp	Leu	Glu	Cys	Ser 320		/ Lys	s Gly	Lys	Cys 325		Thr	Lys	Pro	Ser 330
Glu	Ala	Thr	Phe	Ser 335		s Thr	Cys	Glu	Glu 340		Туг	· Val	. Gly	Thr 345
Phe	Cys	Glu	Glu	350) Ala	a Cys	Gln	355		Pro	су Сув	s Glr	360
Asn	Ala	Ser	Cys	365) Ala	a Asr	ı Glu	1 Lys 370		n Asp	Gly	y Ser	375
Phe	Thr	Cys	; Val	Cys 380		ı Pro	o Gly	7 Tyr	Thr 385		/ Glu	ı Let	ı Cys	390
Ser	Lys	: Ile	e Asp	Ty:		s Ile	e Leu	ı Asp	Pro 400		s Arg	g Ası	n Gly	/ Ala 405





Thr Cys Ile	Ser Ser 410		Ser	Gly	Phe	Thr 415	Cys	Gln	Cys	Pro	Glu 420
Gly Tyr Phe	Gly Ser		Cys	Glu	Glu	Lys 430	Val	Asp	Pro	Cys	Ala 435
Ser Ser Pro	Cys Glr		Asn	Gly	Thr	Cys 445	Tyr	Val	Asp	Gly	Val 450
His Phe Thr	Cys Asr 455		Ser	Pro	Gly	Phe 460	Thr	Gly	Pro	Thr	Cys 465
Ala Gln Leu	Ile Asp		Cys	Ala	Leu	Ser 475	Pro	Суѕ	Ala	His	Gly 480
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Gly Tyr His	Gly Let 50		Cys	Glu	Glu	Glu 505	Tyr	Asn	Glu	Cys	Leu 510
Ser Ala Pro	Cys Let		Ala	Ala	Thr	Cys 520	Arg	Asp	Leu	Val	Asn 525
Gly Tyr Glu	Cys Va	_	Leu	Ala	Glu	Tyr 535	Lys	Gly	Thr	His	Cys 540
Glu Leu Tyr	Lys As		Cys	Ala	Asn	Val 550	Ser	Сув	Leu	Asn	Gly 555
Ala Thr Cys	Asp Se 56		Gly	Leu	Asn	Gly 565	Thr	Cys	Ile	Cys	Ala 570
Pro Gly Phe	Thr Gl 57	-	Glu	. Cys	Asp	Ile 580	Asp	Ile	Asn	Glu	Cys 585
Asp Ser Asn	Pro Cy 59		His	Gly	Gly	Ser 595	Cys	Leu	Asp	Gln	Pro 600
Asn Gly Tyr	Asn Cy		: Cys	Pro	His	Gly 610	Trp	Val	Gly	Ala	Asn 615
Cys Glu Ile	His Le		Trp	Lys	Ser	Gly 625	His	Met	Ala	Glu	Ser 630
Leu Thr Asr	Met Pr 63		His	s Ser	Leu	Tyr 640		Ile	Ile	Gly	Ala 645
Leu Cys Val	Ala Ph		e Leu	: Met	Leu	1le 655		Leu	Ile	· Val	Gly 660
Ile Cys Arg	J Ile Se 66		j Il∈	e Glu	Tyr	670		Ser	Ser	Arg	Pro 675
Ala Tyr Glu	ı Glu Ph 68		. Asr	а Суз	arg	Ser 685		Asp	Ser	Glu	Phe 690
Ser Asn Ala	a Ile Al	a Ser	: Ile	e Arg	, His	Ala	Arg	Ph∈	Gly	/ Lys	Lys

695 700 705

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725 730 735

Asp Leu

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tgtgaggagc agtacgtggg tactttctgt gaagaatacg atgcttgcca 350

gaggaaacct tgccaaaaca acgcgagctg tattgatgca aatgaaaagc 400

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 tctqtqacta agtctattgt ggctttgcgc ttaactctgg tggtgaaggt 200
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Val Tyr Gln Lys Gly Leu Gln Asp Val Asn Leu Arg Asn Phe Ser
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Tyr Gly Gln Thr Ser Leu Asp Arg Leu Arg Asp Gly Leu Val Gly
65 70 75

Ala Gln Phe Trp Ser Ala Tyr Val Pro Cys Gln Thr Gln Asp Arg 80 85 90

Asp Ala Leu Arg Leu Thr Leu Glu Gln Ile Asp Leu Ile Arg Arg 95 100 105

Met Cys Ala Ser Tyr Ser Glu Leu Glu Leu Val Thr Ser Ala Lys 110 115 120

Ala Leu Asn Asp Thr Gln Lys Leu Ala Cys Leu Ile Gly Val Glu 125 130 135

Gly Gly His Ser Leu Asp Asn Ser Leu Ser Ile Leu Arg Thr Phe 140 145 150

Tyr Met Leu Gly Val Arg Tyr Leu Thr Leu Thr His Thr Cys Asn 155 160 165

Thr Pro Trp Ala Glu Ser Ser Ala Lys Gly Val His Ser Phe Tyr 170 175 180



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Asp Ala	Val	Ala	Arg 215	Arg	Ala	Leu	Glu	Val 220	Ser	Gln	Ala	Pro	Val 225
Ile Phe	Ser	His	Ser 230	Ala	Ala	Arg	Gly	Val 235	Cys	Asn	Ser	Ala	Arg 240
Asn Val	Pro	Asp	Asp 245	Ile	Leu	Gln	Leu	Leu 250	Lys	Lys	Asn	Gly	Gly 255
Val Val	Met	Val	Ser 260	Leu	Ser	Met	Gly	Val 265	Ile	Gln	Cys	Asn	Pro 270
Ser Ala	Asn	Val	Ser 275	Thr	Val	Ala	Asp	His 280	Phe	Asp	His	Ile	Lys 285
Ala Val	Ile	Gly	Ser 290	Lys	Phe	Ile	Gly	Ile 295	Gly	Gly	Asp	Tyr	Asp 300
Gly Ala	Gly	Lys	Phe 305	Pro	Gln	Gly	Leu	Glu 310	Asp	Val	Ser	Thr	Tyr 315
Pro Val	Leu	Ile	Glu 320	Glu	Leu	Leu	Ser	Arg 325	Gly	Trp	Ser	Glu	Glu 330
Glu Leu	Gln	Gly	Val 335	Leu	Arg	Gly	Asn	Leu 340	Leu	Arg	Val	Phe	Arg 345
Gln Val	Glu	Lys	Val 350		Glu	Glu	Asn	Lys 355	Trp	Gln	Ser	Pro	Leu 360
Glu Asp	Lys	Phe	Pro 365		Glu	Gln	Leu	Ser 370	Ser	Ser	Cys	His	Ser 375
Asp Leu	Ser	Arg	Leu 380	_	Gln	Arg	Gln	Ser 385		Thr	Ser	Gly	Gln 390
Glu Leu	Thr	Glu	Ile 395		Ile	His	Trp	Thr 400		Lys	Lev	Pro	Ala 405
Lys Trp	Ser	· Val	Ser 410		Ser	Ser	Pro	His 415		Ala	Pro	o Val	Leu 420
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   <211> 50
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    geoctgatge gggactteec getegtggae ggecaeaacg acetgecect 200
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Õ,

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ggcgcccagt tctggtcagc ctatgtgcca tgccagaccc aggaccggga 350

tgccctgcgc ctcaccctgg agcagattga cctcatacgc cgcatgtgtg 400



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Pro Leu Val Asp Gly His Asn Asp Leu Pro Leu Val Leu Arg Gln 35 40 45

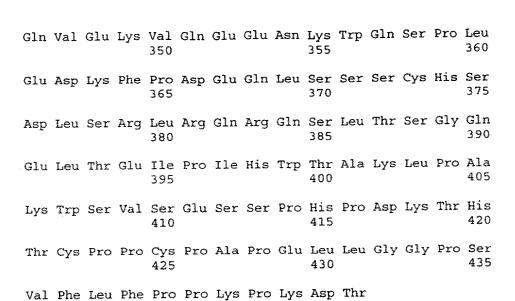
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Asp	Ala	Leu	Arg	Leu 95	Thr	Leu	Glu	Gln	Ile 100	Asp	Leu	Ile	Arg	Arg 105
Met	Cys	Ala	Ser	Tyr 110	Ser	Glu	Leu	Glu	Leu 115	Val	Thr	Ser	Ala	Lys 120
Ala	Leu	Asn	Asp	Thr 125	Gln	Lys	Leu	Ala	Cys 130	Leu	Ile	Gly	Val	Glu 135
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Tyr	Met	Leu	Gly	Val 155	Arg	Tyr	Leu	Thr	Leu 160	Thr	His	Thr	Cys	Asn 165
Thr	Pro	Trp	Ala	Glu 170	Ser	Ser	Ala	Lys	Gly 175	Val	His	Ser	Phe	Tyr 180
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Glu	Met	Asn	Arg	Leu 200	Gly	Met	Met	Val	Asp 205	Leu	Ser	His	Val	Ser 210
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Ile	Phe	Ser	His	Ser 230		Ala	Arg	Gly	Val 235	Cys	Asn	Ser	Ala	Arg 240
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Gly	Ala	Gly	Lys	Phe		Glr	Gly	Leu	Glu 310		Val	Ser	Thr	Tyr 315
Pro	Val	Leu	Ile	Glu 320		Let	ı Leu	Ser	Arg 325		Trp	Ser	Glu	330
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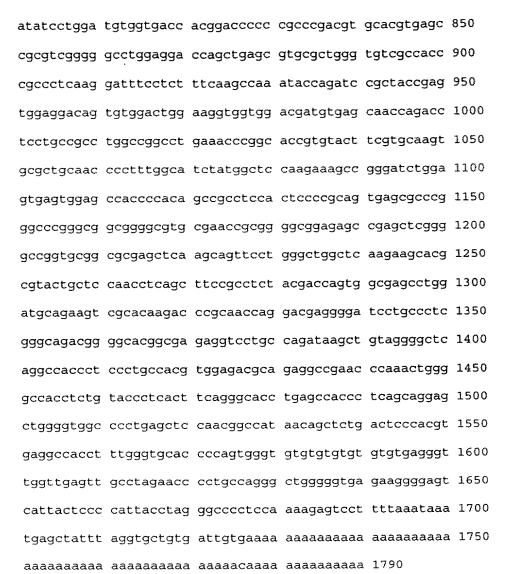


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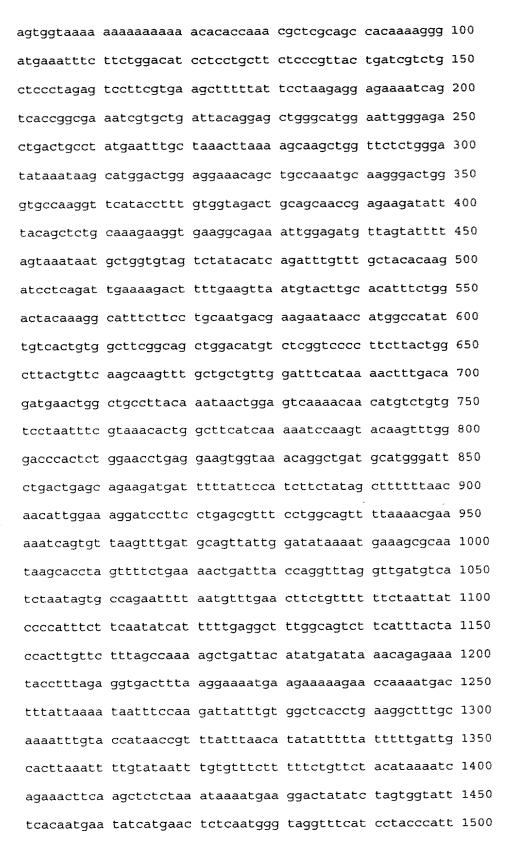




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His Ile Pro	Lys Asp 200	Leu	Ala	Leu	Phe	Thr 205	Pro	Tyr	Glu	Ile	Trp 210
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Thr Leu Asp	Ile Leu 230		Val	Val	Thr	Thr 235	Asp	Pro	Pro	Pro	Asp 240
Val His Val	Ser Arg 245		Gly	G1y	Leu	Glu 250	Asp	Gln	Leu	Ser	Val 255
Arg Trp Val	Ser Pro		Ala	Leu	Lys	Asp 265	Phe	Leu	Phe	Gln	Ala 270
Lys Tyr Gln	Ile Arg 275	_	Arg	Val	Glu	Asp 280	Ser	Val	Asp	Trp	Lys 285
Val Val Asp	Asp Val		Asn	Gln	Thr	Ser 295	Cys	Arg	Leu	Ala	Gly 300
Leu Lys Pro	Gly Thr		Tyr	Phe	Val	Gln 310	Val	Arg	Cys	Asn	Pro 315
Phe Gly Ile	Tyr Gly		Lys	Lys	Ala	Gly 325	Ile	Trp	Ser	Glu	330
Ser His Pro	Thr Ala		Ser	Thr	Pro	Arg 340	Ser	Glu	Arg	Pro	Gly 345
Pro Gly Gly	Gly Ala	Cys	Glu	Pro	Arg	Gly	Gly	Glu	Pro	Ser	Ser

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35 40 45

His Gly Ile Gly Arg Leu Thr Ala Tyr Glu Phe Ala Lys Leu Lys
50 55 60

Ser Lys Leu Val Leu Trp Asp Ile Asn Lys His Gly Leu Glu Glu 65 70 75

Thr Ala Ala Lys Cys Lys Gly Leu Gly Ala Lys Val His Thr Phe 80 85 90

Val Val Asp Cys Ser Asn Arg Glu Asp Ile Tyr Ser Ser Ala Lys 95 100 105

Lys Val Lys Ala Glu Ile Gly Asp Val Ser Ile Leu Val Asn Asn 110 115 120

Ala Gly Val Val Tyr Thr Ser Asp Leu Phe Ala Thr Gln Asp Pro 125 130 135

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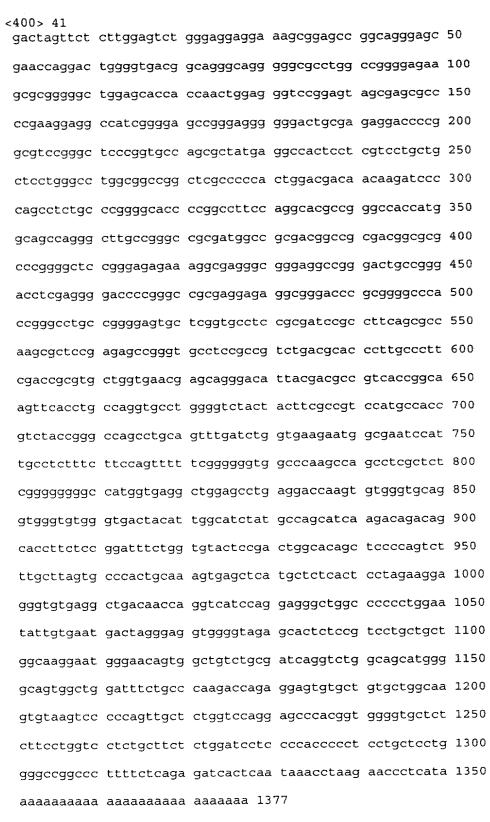
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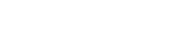
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Ala Pro Gly Glu Lys Gly Glu Gly Gly Arg Pro Gly Leu Pro Gly
65 70 75

Pro Arg Gly Asp Pro Gly Pro Arg Gly Glu Ala Gly Pro Ala Gly 80 85 90

Pro Thr Gly Pro Ala Gly Glu Cys Ser Val Pro Pro Arg Ser Ala 95 100 105

Phe Ser Ala Lys Arg Ser Glu Ser Arg Val Pro Pro Pro Ser Asp 110 115 120

Ala Pro Leu Pro Phe Asp Arg Val Leu Val Asn Glu Gln Gly His
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Tyr Asp Ala Val Thr Gly Lys Phe Thr Cys Gln Val Pro Gly Val

Tyr Tyr Phe Ala Val His Ala Thr Val Tyr Arg Ala Ser Leu Gln
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Phe Asp Leu Val Lys Asn Gly Glu Ser Ile Ala Ser Phe Phe Gln
170 175 180

Phe Phe Gly Gly Trp Pro Lys Pro Ala Ser Leu Ser Gly Gly Ala 185 190 195

Met Val Arg Leu Glu Pro Glu Asp Gln Val Trp Val Gln Val Gly
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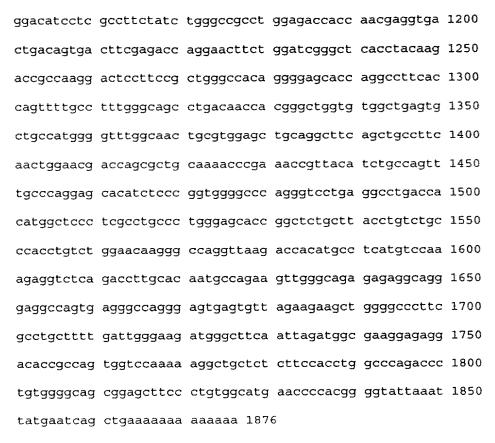
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Pro Gln Leu Gln Glu Gln Ala Pro Met Ala Gly Ala Leu Asn Arg 35 40 45

Lys Glu Ser Phe Leu Leu Leu Ser Leu His Asn Arg Leu Arg Ser 50 55 60

Trp Val Gln Pro Pro Ala Ala Asp Met Arg Arg Leu Asp Trp Ser
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Asp Ser Leu Ala Gln Leu Ala Gln Ala Arg Ala Ala Leu Cys Gly 80 85 90

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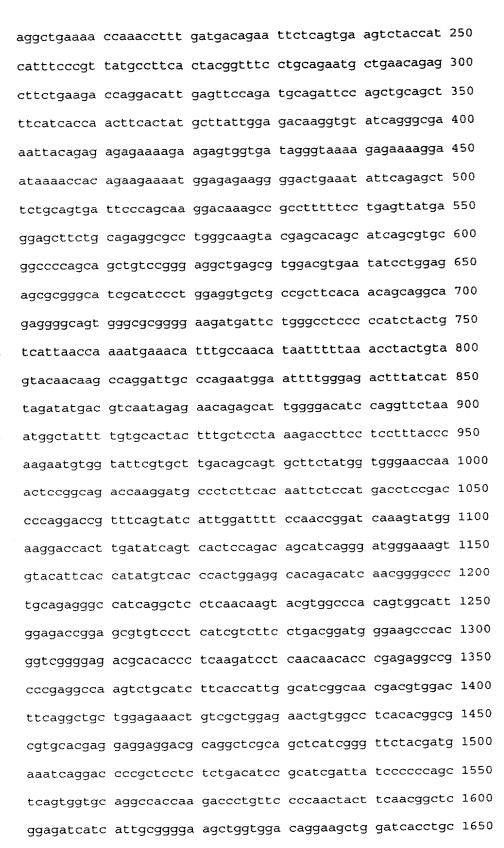




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His Leu	Cys	Ser	Ala 170	Gly	Gln	Thr	Ala	Ile 175	Glu	Ala	Phe	Val	Cys 180
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Tyr Gly	y Gly	Ala	Gln 290	Cys	Ala	Thr	Lys	Val 295		Phe	Pro	Phe	His 300
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Gly Gl	y Val	Leu	Ala 335		Ile	Lys	Ser	Gln 340		Val	Gln	Asp	345
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Asp Se	r Asp	Phe	Glu 365		Arg	Asn	Phe	Trp		e Gly	Leu	Thr	Tyr 375
Lys Th	r Ala	Lys	Asp 380		Phe	Arg	Trp	385		Gly	/ Glu	ı His	390
Ala Ph	e Thi	Ser	Phe 395		Phe	Gly	Glr	400		Asr	ı His	Gl _y	/ Leu 405

Val Trp Leu Ser Ala Ala Met Gly Phe Gly Asn Cys Val Glu Leu 410 415 Gln Ala Ser Ala Ala Phe Asn Trp Asn Asp Gln Arg Cys Lys Thr Arg Asn Arg Tyr Ile Cys Gln Phe Ala Gln Glu His Ile Ser Arg Trp Gly Pro Gly Ser 455 <210> 51 <211> 24 <212> DNA <213> Artificial Sequence <220> <223> Synthetic oligonucleotide probe <400> 51 aggaacttct ggatcgggct cacc 24 <210> 52 <211> 24 <212> DNA <213> Artificial Sequence <223> Synthetic oligonucleotide probe <400> 52 gggtctgggc caggtggaag agag 24 <210> 53 <211> 45 <212> DNA <213> Artificial Sequence <220> <223> Synthetic oligonucleotide probe gecaaggact cetteegetg ggccaeaggg gagcaeeagg cette 45 <210> 54 <211> 2331 <212> DNA <213> Homo Sapien <400> 54 cggacgcgtg ggctgggcgc tgcaaagcgt gtcccgccgg gtccccgagc 50 gtcccgcgcc ctcgccccgc catgctcctg ctgctggggc tgtgcctggg 100 gctgtccctg tgtgtggggt cgcaggaaga ggcgcagagc tggggccact 150 cttcggagca ggatggactc agggtcccga ggcaagtcag actgttgcag 200

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35 40 45

Lys Thr Lys Pro Leu Met Thr Glu Phe Ser Val Lys Ser Thr Ile
50 55 60

Ile Ser Arg Tyr Ala Phe Thr Thr Val Ser Cys Arg Met Leu Asn 65 70 75

Arg Ala Ser Glu Asp Gln Asp Ile Glu Phe Gln Met Gln Ile Pro
80 85 90

Ala Ala Ala Phe Ile Thr Asn Phe Thr Met Leu Ile Gly Asp Lys 95 100 105

Val Tyr Gln Gly Glu Ile Thr Glu Arg Glu Lys Lys Ser Gly Asp 110 115 120



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Asp	Lys	Ala	Ala	Phe 155	Phe	Leu	Ser	Tyr	Glu 160	Glu	Leu	Leu	Gln	Arg 165
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Leu	Ser	Gly	Arg	Leu 185	Ser	Val	Asp	Val	Asn 190	Ile	Leu	Glu	Ser	Ala 195
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Tyr	Ile	e His	s His	365		Pro	Thr	Gly	Gly 370		Asp) Ile	e Asn	375
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Ser	Gly	/ Ile	e Gly	/ Asp 395		Sei	c Val	. Ser	Let 400		e Val	l Ph∈	e Lev	1 Thr 405
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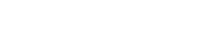


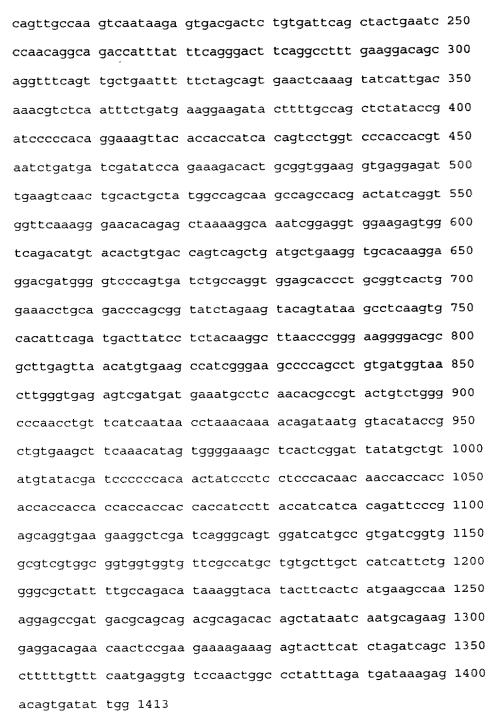


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Pro	Gln	Pro	Val	Met 275	Val	Thr	Trp	Val	Arg 280	Val	Asp	Asp	Glu	Met 285
Pro	Gln	His	Ala	Val 290	Leu	Ser	Gly	Pro	Asn 295	Leu	Phe	Ile	Asn	Asn 300
Leu	Asn	Lys	Thr	Asp	Asn	Gly	Thr	Tyr	Arg	Cys	Glu	Ala	Ser	Asn



315 305 310 Ile Val Gly Lys Ala His Ser Asp Tyr Met Leu Tyr Val Tyr Asp Pro Pro Thr Thr Ile Pro Pro Pro Thr Thr Thr Thr Thr Thr 335 Thr Thr Thr Thr Thr Ile Leu Thr Ile Ile Thr Asp Ser Arg 350 355 Ala Gly Glu Glu Gly Ser Ile Arg Ala Val Asp His Ala Val Ile 365 370 Gly Gly Val Val Ala Val Val Phe Ala Met Leu Cys Leu Leu Ile Ile Leu Gly Arg Tyr Phe Ala Arg His Lys Gly Thr Tyr Phe Thr His Glu Ala Lys Gly Ala Asp Asp Ala Ala Asp Ala Asp Thr Ala Ile Ile Asn Ala Glu Gly Gly Gln Asn Asn Ser Glu Glu Lys Lys Glu Tyr Phe Ile <210> 62 <211> 24 <212> DNA <213> Artificial Sequence <223> Synthetic oligonucleotide probe <400> 62 ggettetget gttgetette teeg 24 <210> 63 <211> 20 <212> DNA <213> Artificial Sequence <220> <223> Synthetic oligonucleotide probe <400> 63 gtacactgtg accagtcagc 20 <210> 64 <211> 20 <212> DNA <213> Artificial Sequence <220>

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Ser Gln Pro Gln Thr Val Phe Cys Thr Ala Arg Gln Gly Thr Thr 35 40 45

Val Pro Arg Asp Val Pro Pro Asp Thr Val Gly Leu Tyr Val Phe
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Glu Asn Gly Ile Thr Met Leu Asp Ala Ser Ser Phe Ala Gly Leu 65 70 75

Pro Gly Leu Gln Leu Leu Asp Leu Ser Gln Asn Gln Ile Ala Ser 80 85 90

Leu Arg Leu Pro Arg Leu Leu Leu Leu Asp Leu Ser His Asn Ser 95 100 105

Leu Leu Ala Leu Glu Pro Gly Ile Leu Asp Thr Ala Asn Val Glu

110 115 120

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Leu	Phe	Ser	Arg	Leu 140	Arg	Asn	Leu	His	Asp 145	Leu	Asp	Val	Ser	Asp 150
Asn	Gln	Leu	Glu	Arg 155	Val	Pro	Pro	Val	Ile 160	Arg	Gly	Leu	Arg	Gly 165
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Arg	Pro	Glu	Asp	Leu 185	Ala	Gly	Leu	Ala	Ala 190	Leu	Gln	Glu	Leu	Asp 195
Val	Ser	Asn	Leu	Ser 200	Leu	Gln	Ala	Leu	Pro 205	Gly	Asp	Leu	Ser	Gly 210
Leu	Phe	Pro	Arg	Leu 215	Arg	Leu	Leu	Ala	Ala 220	Ala	Arg	Asn	Pro	Phe 225
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Ser	His	Val	Thr	Leu 245	Ala	Ser	Pro	Glu	Glu 250	Thr	Arg	Сув	His	Phe 255
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<223> Synthetic oligonucleotide probe

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<213> Homo Sapien

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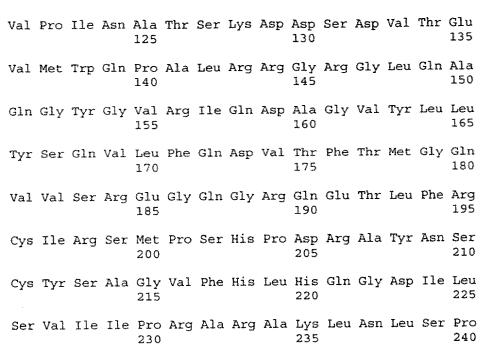
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Glu Val Ser Arg Leu Gln Gly Thr Gly Gly Pro Ser Gln Asn Gly 65 70 75

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Leu Glu Ala Trp Glu Asn Gly Glu Arg Ser Arg Lys Arg Ala 95 100 105

Val Leu Thr Gln Lys Gln Lys Gln His Ser Val Leu His Leu 110 $\,$ 115 $\,$ 120



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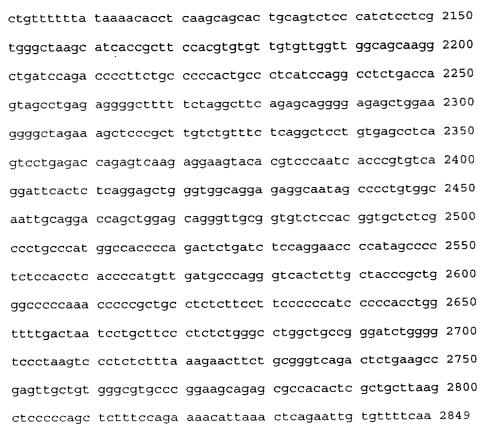
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<212> PRT

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Gln Gly Glu Gln Gln Glu Trp Glu Gly Thr Glu Glu Leu Pro Ser 35 40 45

Pro Pro Asp His Ala Glu Arg Ala Glu Glu Gln His Glu Lys Tyr
50 55 60

Arg Pro Ser Gln Asp Gln Gly Leu Pro Ala Ser Arg Cys Leu Arg
65 70 75

Cys Cys Asp Pro Gly Thr Ser Met Tyr Pro Ala Thr Ala Val Pro 80 85 90

Gln Ile Asn Ile Thr Ile Leu Lys Gly Glu Lys Gly Asp Arg Gly 95 100 105

Asp Arg Gly Leu Gln Gly Lys Tyr Gly Lys Thr Gly Ser Ala Gly

115 120 110 Ala Arg Gly His Thr Gly Pro Lys Gly Gln Lys Gly Ser Met Gly Ala Pro Gly Glu Arg Cys Lys Ser His Tyr Ala Ala Phe Ser Val Gly Arg Lys Lys Pro Met His Ser Asn His Tyr Tyr Gln Thr Val 155 Ile Phe Asp Thr Glu Phe Val Asn Leu Tyr Asp His Phe Asn Met 170 Phe Thr Gly Lys Phe Tyr Cys Tyr Val Pro Gly Leu Tyr Phe Phe Ser Leu Asn Val His Thr Trp Asn Gln Lys Glu Thr Tyr Leu His 200 Ile Met Lys Asn Glu Glu Glu Val Val Ile Leu Phe Ala Gln Val Gly Asp Arg Ser Ile Met Gln Ser Gln Ser Leu Met Leu Glu Leu Arg Glu Gln Asp Gln Val Trp Val Arg Leu Tyr Lys Gly Glu Arg Glu Asn Ala Ile Phe Ser Glu Glu Leu Asp Thr Tyr Ile Thr Phe Ser Gly Tyr Leu Val Lys His Ala Thr Glu Pro 275 <210> 79 <211> 24 <212> DNA <213> Artificial Sequence <223> Synthetic oligonucleotide probe <400> 79 tacaggccca gtcaggacca gggg 24 <210> 80 <211> 24 <212> DNA <213> Artificial Sequence <220> <223> Synthetic oligonucleotide probe <400> 80 ctgaagaagt agaggccggg cacg 24 <210> 81



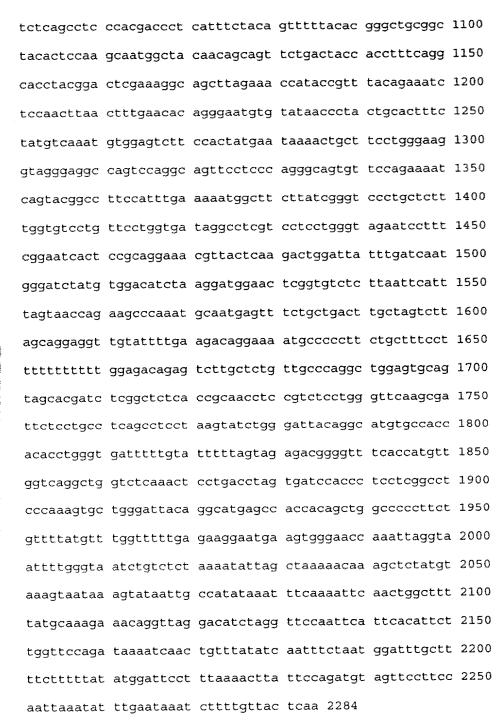


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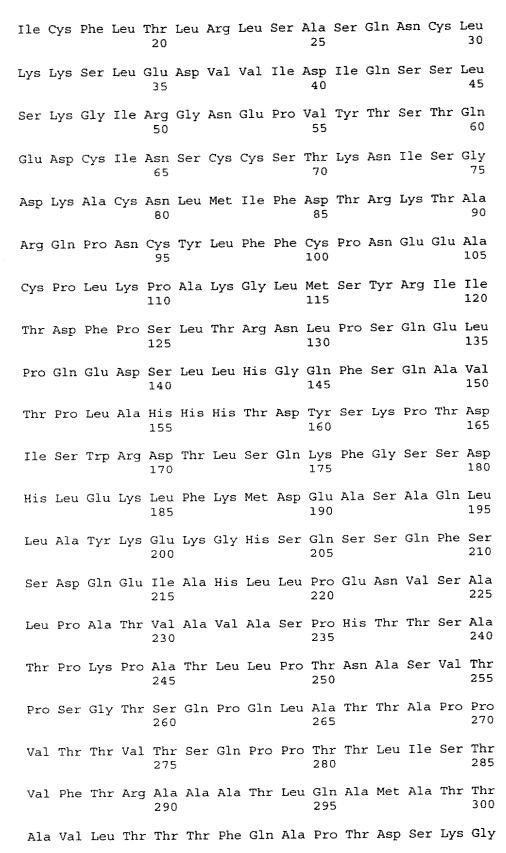
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Val Glu Ser Ser	Thr 350	Met	Asn	Lys	Thr	Ala 355	Ser	Trp	Glu	Gly	Arg 360	
Glu Ala Ser Pro	Gly 365	Ser	Ser	Ser	Gln	Gly 370	Ser	Val	Pro	Glu	Asn 375	
Gln Tyr Gly Leu	Pro 380	Phe	Glu	Lys	Trp	Leu 385	Leu	Ile	Gly	Ser	Leu 390	
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35 40 45

Ser Ile Gly Glu Arg Pro Val Leu Lys Ala Pro Val Pro Lys Arg
50 55 60

Gln Lys Cys Asp His Trp Thr Pro Cys Pro Ser Asp Thr Tyr Ala 65 70 75

Tyr Arg Leu Leu Ser Gly Gly Gly Arg Ser Lys Tyr Ala Lys Ile 80 85 90

Cys Phe Glu Asp Asn Leu Leu Met Gly Glu Gln Leu Gly Asn Val 95 100 105

Ala Arg Gly Ile Asn Ile Ala Ile Val Asn Tyr Val Thr Gly Asn 110 115 120

Val Thr Ala Thr Arg Cys Phe Asp Met Tyr Glu Gly Asp Asn Ser 125 130 135

Gly Pro Met Thr Lys Phe Ile Gln Ser Ala Ala Pro Lys Ser Leu 140 145 150

Leu Phe Met Val Thr Tyr Asp Asp Gly Ser Thr Arg Leu Asn Asn 155 160 165

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